



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/700,700

DATE: 08/13/2002 TIME: 14:16:34

Input Set : A:\dex-105.ST25.txt

Output Set: N:\CRF4\08132002\I700700.raw

ENTERED

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3 <110> APPLICANT: Ali, Shujath
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- 4 Salceda, Susana
- 5 Sun, Yongming
- 6 Cafferkey, Robert
- 8 <120> TITLE OF INVENTION: A NOVEL METHOD OF DIAGNOSING, MONITORING, AND STAGING PROSTATE CANCER
 - 10 <130> FILE REFERENCE: DEX-0105
 - 12 <140> CURRENT APPLICATION NUMBER: 09/700,700
 - 13 <141> CURRENT FILING DATE: 2000-11-20
 - 15 <150> PRIOR APPLICATION NUMBER: PCT/US99/10548
 - 16 <151> PRIOR FILING DATE: 1999-05-12
 - 18 <150> PRIOR APPLICATION NUMBER: US 60/086,265
 - 19 <151> PRIOR FILING DATE: 1998-05-21
 - 21 <160> NUMBER OF SEQ ID NOS: 7
 - 23 <170> SOFTWARE: PatentIn version 3.1
 - 25 <210> SEQ ID NO: 1
 - 26 <211> LENGTH: 1936
 - 27 <212> TYPE: DNA
 - 28 <213> ORGANISM: Homo sapiens
 - 30 <220> FEATURE:
 - 31 <221> NAME/KEY: misc_feature
 - 32 <222> LOCATION: (106)..(106)
 - 33 <223> OTHER INFORMATION: n=a, c, g or t
 - 36 <220> FEATURE:
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 - 57 <223> OTHER INFORMATION: n=a, c, g or t
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 - 61 <221> NAME/KEY: misc_feature
 - 62 <222> LOCATION: (1908)..(1908)
 - 63 <223> OTHER INFORMATION: n=a, c, g or t
 - 66 <400> SEQUENCE: 1

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Output Set: N:\CRF4\08132002\1700700.raw

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M>	> 69 tgattttaag acaagctgta taaacagaac tccactgcaa gagggnggg		120
	71 gaatctccgc ttgtccaaga caggggccta aggagggtct ccacactgc		180
	73 gttgcatttt tttattagta gaaagtggaa aggcctcttc tcaactttt		240
	75 tggagaattt agaatcagaa gtttcctgga gttttcaggc tatcatata		300
	77 gaaaggcaac ataattette etteceteet tttaaaattt tgtgtteet		360
	79 ttactcacta aagggcttca ttttagtcca gatttttagt ctggctgca	c ctaacttatg	420
M>	81 cctcgcttat ttagcccgag atctggtctt ttttntgtnt ttttttnt		480
	83 caaagcttta totgtottga otttttaaaa aagtttgggg goagattot		540
	85 aaagacatgc atttttaaaa ctaggcaact tcttatttct ttcctttaa	a aatacatagc	600
	87 attaaatccc aaatcctatt taaagacctg acagcttgag aaggtcact		660
	89 aggacettet ggtggttetg etgttacgtt tgaagtetga caateettg	a gaatctttgc	720
	91 atgcagagga ggtaagaggt attggatttt cacagaggaa gaacacagc	g cagaatgaag	780
	93 ggccaggett actgaggetg tecagtggag ggctcatggg tgggacatg	g aaaagaaggc	840
	95 agectaggee etggggagee cagtecactg ageaageaag ggactgagte		900
	97 aggaaaaggc taagaaaaag gaaaaccatt ctaaaacaca acaagaaac	t gtccaaatgc	960
	99 tttgggaact gtgtttattg cctataatgg gtccccaaaa tgggtaacc		1020
	101 gagaatgagc agagagcaaa ggagaaatct ggctgtcctt ccattttc		1080
	103 caggtgagct ggtagagggg agacattaga aaaaaatgaa acaacaaa		1140
	105 gaggtacgct gaggcctggg agtctcttga ctccactact taattccg		1200
	107 cctttcaatt ttcttttatt agaagggcca gcttactgtt ggtggcaa		1260
	109 aagttaatag aaagttggcc aatttcaccc cattttctgt ggtttggg		1320
	111 atgttcaatg ccacgtgctg ctgacaccga ccggagtact agccagca		1380
	113 tagcctgaat tgctttctgc tctttacatt tcttttaaaa taagcatt		1440
	115 cctactgagt actctttctc tcccctcctc tgaatttaat tctttcaa		1500
W>	> 117 aaggattaca catttcactg tgatgtatat tgtgttgcag ngaaaaga		1560
	119 tgtttaaaat tacttggttt gtgaatccat cttgcttttt ccccattg		1620
	121 aacccatctc tgaactggta gaaaaacatc tgaagagcta gtctatca		1680
	123 tgaattggat ggttctcaga accatttcac ccagacagcc tgtttcta		1740
	125 aattagtttg ggttctctac atgcataaca aaccctgctc caatctgt		1800
	127 tgtgacttga agtttagtca gcacccccac caaactttat ttttctat		1860
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	135 <211> LENGTH: 637		
	136 <212> TYPE: DNA		
	137 <213> ORGANISM: Homo sapiens		
	139 <400> SEQUENCE: 2		
	140 gtaggggcag acttactgcc ttgaacgaaa gacgatggtc ctcgctca	re etcaetecaa	60
	142 ttatgtteet etaggtgggg caggtagggg gtecagette etgettge		120
	144 teatgestee ageettstee ettetsacet gggeettee eagggga		180
	146 gcagaagaat cagccccaca gtgcaggggt gtgttagtgg ggaacggg		240
			300
	148 gtgggaacca gggaccccct atcttggtac cggtcattgg atgtatcc 150 tgtgtctgtc ttggcccgtg tggtcaccct gtgttcatct ctctcca		360
			420
	152 caaactgggg ttttcgtctc cctatgaggg ggtcctggta tgtacgcg		420
	154 geggtgeatg teteceggtg cagtgeatge tggggtteee tggggeee		540
	156 aggatagaca gagcotgtoc taacottoog gaagtgcatg otggggag		600
	158 ctgaccttct gtgctcagga cgactaatcg gccacatgac caccactc 160 ttcctagaga agtctcacta agagcccagc acactca	ig coccat <u>y</u> gga	637
	100 cccccayaya aycoccacca ayayoocayo acaccca		557

Input Set : A:\dex-105.ST25.txt

Output Set: N:\CRF4\08132002\I700700.raw

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     171 <223> OTHER INFORMATION: n=a, c, g or t
     174 <220> FEATURE:
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     176 <222> LOCATION: (1480)..(1480)
     177 <223> OTHER INFORMATION: n=a, c, g or t
     180 <220> FEATURE:
     181 <221> NAME/KEY: misc_feature
     182 <222> LOCATION: (1532)..(1532)
     183 <223> OTHER INFORMATION: n=a, c, g or t
     186 <220> FEATURE:
     187 <221> NAME/KEY: misc_feature
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     189 <223> OTHER INFORMATION: n=a, c, g or t
     192 <220> FEATURE:
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     194 <222> LOCATION: (1569)..(1569)
     195 <223> OTHER INFORMATION: n=a, c, g or t
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     199 <221> NAME/KEY: misc_feature
     200 <222> LOCATION: (1571)..(1571)
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     206 <222> LOCATION: (1631)..(1631)
     207 <223> OTHER INFORMATION: n=a, c, g or t
     210 <220> FEATURE:
     211 <221> NAME/KEY: misc_feature
     212 <222> LOCATION: (2266)..(2512)
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     219 ccacagccat cctaggagaa aggcagaatg tcataatttg ctaaaaggga tgctgaggct
                                                                               120
     221 ctgggaggga aagggacttg cctaaagccc cagggtgaag cagcatctct ggactcccag
                                                                               180
     223 tecagtgate ttgeecaata etttgetget tgeetatace cetetaactt ggteaacage
                                                                               240
     225 acatcacagg gcaagcccaa tccctgcttc atttttatat atgggcgctg gtccacagcc
                                                                               300
     227 ccactctcca gccatttgga aacaaaaaca gatgctattg ttcttcctta gagaacgtgg
     229 ccagtggaga cggcacactg gaaatcagag tgaatgttct tgaaagaggg tcacgggtca
                                                                               420
     231 acaaggccca gccaaaggat gcagtagaac cattttcctt agaaatcttt gggagtgaag
                                                                               480
     233 taggetteag ceactaceea tecetgeeet tgeggetace actaceceat tagtttagae
                                                                               540
W--> 235 agggtcgggc ggggaggggt gtggagaaga aatgagcttg cctgtngccc ccaggctccc
                                                                               600
     237 tetgteetag eteaggtetg ggtgeeatte tttacaeteg tgtgeteget caegeacaea
                                                                               660
     239 tcacacacct tgctggtcac acagtcacag actcgcctct gctcctgtgg tccagtggcc
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Input Set : A:\dex-105.ST25.txt

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    243 tgaaggaaat ccacacaccc agagcatctc ggagttcaga ctctcagacc tgaagtaggc
                                                                        840
                                                                       900
    245 gcccccggga ctgggctagg agttggacgg aatggaggat ggaggacagc gagaagaaag
    247 gaagagaaat gcaaagtgtg ggcagccgcc aagagtgaaa atagagggaa gtgtcatgca
                                                                       960
    249 agtgctggac agaaggcggc aggtgggacg agccccacag cccctcctc aaaaacgacc
                                                                       1020
    251 acctccagga ctcagtgatc cctggggggc aggctctgcc agccctcggc cacacgtggc
                                                                       1080
                                                                       1140
    253 teeggeacee atggteecag tgeettggat ggagaeggee agttetggeg geeagatgtg
    255 gtgctctgga atccagtccc atttccttcc tggccacgcc tgttccagcg gcctctttgg
                                                                       1200
    257 ctgcattcag cccctactta cctggggacc ccggctgggg cacaagagca ccaggggggt
                                                                       1260
    259 agggcccaaa gggatcaggg gaagcctctg gcctggaggg tatggggcac gcttccccaa
                                                                       1320
    261 gggcggaccc ggcaggagga agcccaggag ctgggtcctg ccgcccagga gctgggccct
                                                                       1380
    263 gccacccagg ccgggctagg gacatggcag ggcctgggca tcctgacgct ggacttgggc
                                                                       1440
                                                                       1500
W--> 265 qacctqqqaq qcacaqqqaq qqqaqaqatq ggcqgccccn acccagcgca gtgccggcca
W--> 267 caccccaagg cggttgccag agcttaaggc cnggccccag caggagaaca tcccagctcc
                                                                       1560
W--> 269 annnnncene neegeageea gtgeteettg teaageteee eeegteacte eaggtgggag
                                                                       1620
W--> 271 ccacccggt nagggggtgt gccacttgcc cccagggcac tcctctgggc atcccgggtg
                                                                       1680
                                                                       1740
    273 ggggattttg gggccgtggg gggcagtctc tggtacctgt gtgcgtcagg gatgctctgc
                                                                       1800
    275 acctgcaacc aggtgtcgtc cacgggcggg ggcatgggca tggtgacagt ggtcctgttg
                                                                       1860
    277 atgtcaccga tgatgctgag cgcctccttc agcgcgtggt gcatgtgcag catctcgtcg
    279 tgctgctgtg cctgctctgc caactcctcc atcagtgtgt tctggttccc acatgagtac
                                                                       1920
    281 atattggcca gcggctccga gatgatgaac tccggggtct gagagtgggc aaacagggaa
                                                                       1980
    283 gaaggttggg acctggtgcc tgtgccgccc tggctgcctt gctgggccct tctgggactg
                                                                       2040
                                                                       2100
    285 tgcgctggac ttggagcccc ttggagtatg gcttttcaca cgggcttcta taccgcttcg
    287 actggaagat ccacctcccc actgcctttt ctcactcaga tggggacacc gaggtccaga
                                                                       2160
                                                                       2220
    289 ggaaaagaca cctgtcaaat gtcacagatc tgggagggga cttaagacct atcatgccaa
                                                                       2280
W--> 291 gaggacacct gtctactcag ttttttttttg gtggggcggg gggcgnnnnn nnnnnnnnn
2340
                                                                       2400
2460
2520
                                                                       2580
    301 agttgatgcc tggatacagg agctctgtgg gtgggagtga gacaaaacac agggtcctga
                                                                       2640
    303 gctctgggga ccaagcaatg tcctctggtg aaaaaaatcc tggacttgct ggcagaagat
    305 ttgcctctta cttgccatgt gctctgaata catttacctg ccctctggga aaa
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                                                                        120
    322 aactatccag taaggcttag agggtttcaa tcagaaatat gtgttagggg aaaaaatgca
    324 ctttttctat attaaaaaat attattttct tcttttaaat gtaaagcatt cctattgtga
                                                                        180
    326 agaattgaga aaatacagaa aagtacaaag aaaaacatta cctacaactc caccatccgt
                                                                        240
                                                                        292
W--> 328 gattatcact gttcacattt gtggctcatt tttcagtatk tctnttattt aa
    331 <210> SEQ ID NO: 5
    332 <211> LENGTH: 2694
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Input Set : A:\dex-105.ST25.txt

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334 <213> ORGANISM: Homo sapiens
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380 <222> LOCATION: (132)..(132)
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392 <222> LOCATION: (217)..(217)
393 <223> OTHER INFORMATION: n=a, c, g or t
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398 <222> LOCATION: (257)..(257)
399 <223> OTHER INFORMATION: n=a, c, g or t
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403 <221> NAME/KEY: misc_feature
404 <222> LOCATION: (2539)..(2539)
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/700,700

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:3; N Pos. 586,1480,1532,1562,1563,1564,1565,1566,1569,1571,1631,2266
Seq#:3; N Pos. 2267,2268,2269,2270,2271,2272,2273,2274,2275,2276,2277,2278
Seq#:3; N Pos. 2279,2280,2281,2282,2283,2284,2285,2286,2287,2288,2289,2290
Seq#:3; N Pos. 2291,2292,2293,2294,2295,2296,2297,2298,2299,2300,2301,2302
Seq#:3; N Pos. 2303,2304,2305,2306,2307,2308,2309,2310,2311,2312,2313,2314
Seq#:3; N Pos. 2315,2316,2317,2318,2319,2320,2321,2322,2323,2324,2325,2326
Seq#:3; N Pos. 2327,2328,2329,2330,2331,2332,2333,2334,2335,2336,2337,2338
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Seq#:3; N Pos. 2351,2352,2353,2354,2355,2356,2357,2358,2359,2360,2361,2362
Seq#:3; N Pos. 2363,2364,2365,2366,2367,2368,2369,2370,2371,2372,2373,2374
Seq#:3; N Pos. 2375,2376,2377,2378,2379,2380,2381,2382,2383,2384,2385,2386
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Seq#:3; N Pos. 2495,2496,2497,2498,2499,2500,2501,2502,2503,2504,2505,2506
Seg#:3; N Pos. 2507,2508,2509,2510,2511,2512
Seg#:4; N Pos. 284 / / / /
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Seq#:6; N Pos. 17
Seq#:7; N Pos. 268,688,700
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